

**AMENDMENTS TO THE CLAIMS**

Claims 1-24 (Cancelled)

25. (Previously presented): A method to identify a compound which behaves as an agonist for a T-type calcium channel which method comprises:

a) contacting a recombinant cell which expresses the  $\alpha_1$  subunit of a heterologous T-type calcium channel with a compound to be tested; and

b) determining the ability of said compound to activate said  $\alpha_1$  subunit;

wherein said  $\alpha_1$  subunit is functional as a T-type calcium ion channel and is encoded by a nucleotide sequence which hybridizes under conditions of stringency corresponding to washing at 62° C in 0.2 x SSPE/0.1% SDS to a nucleic acid comprising SEQ ID NO: 23, 25 or 27, and

wherein said activating comprises enhancing the flow of calcium ions into said cell in the presence as compared to the absence of said compound;

whereby a compound which activates said  $\alpha_1$  subunit is identified as an agonist of said T-type calcium channel.

26. (Previously presented): The method of claim 25 wherein said activation is measured by measuring the current through the calcium channel before and after said contacting of said cell with said compound.

27. (Previously presented): The method of claim 25, wherein said cells contain a fluorescent dye sensitive to intracellular calcium concentration and said activation is determined by observing a change in the fluorescence of said dye when said contacting is performed.

28. (Previously presented): A method to identify an antagonist of a T-type calcium channel which method comprises:

a) contacting a recombinant cell expressing the  $\alpha_1$  subunit of a heterologous T-type calcium channel with a known agonist of said T-type calcium channel;

b) contacting said cell with a compound to be tested; and

c) determining the ability of said compound to diminish the activation of said  $\alpha_1$  subunit by said agonist;

wherein said  $\alpha_1$  subunit is functional as a T-type calcium ion channel and is encoded by a nucleotide sequence which hybridizes under conditions of stringency corresponding to washing at 62° C in 0.2 x SSPE/0.1% SDS to a nucleic acid comprising SEQ ID NO: 23, 25 or 27, and

wherein said activating comprises enhancing the flow of calcium ions into said cell in the presence as compared to the absence of said agonist;

whereby a compound which diminishes the activation of said  $\alpha_1$  subunit by said agonist is identified as an antagonist.

29. (Previously presented): The method of claim 28 wherein said activation is measured by measuring the current through the calcium channel before and after said contacting of said cell with said compound.

30. (Previously presented): The method of claim 28, wherein said cells contain a fluorescent dye sensitive to intracellular calcium concentration and said activation is determined by observing a change in the fluorescence of said dye when said contacting is performed.

31. (Previously presented): A method to prescreen compounds as agonists or antagonists of T-type calcium ion channels by virtue of their ability to bind said T-type channels which method comprises:

a) contacting a recombinant cell expressing the  $\alpha_1$  subunit of a heterologous T-type calcium channel with a compound to be tested; and

b) determining the ability of said compound to bind to said cell expressing said  $\alpha_1$  subunit;

wherein said binding is determined by observing competitive binding with a known agonist or antagonist of said channel;

wherein said  $\alpha_1$  subunit is functional as a T-type calcium ion channel and is encoded by a nucleotide sequence which hybridizes under conditions of stringency corresponding to washing at 62°C in 0.2 x SSPE/0.1% SDS to a nucleic acid comprising SEQ ID NO: 23, 25 or 27,

whereby a compound which is determined to bind said cell is identified as a compound which will behave as either an agonist or antagonist of a T-type calcium channel.

32. (New): The method defined in claim 25 wherein the nucleic acid comprises SEQ ID NO: 23.

33. (New): The method defined in claim 25 wherein the nucleic acid comprises SEQ ID NO: 25.

34. (New): The method defined in claim 25 wherein the nucleic acid comprises SEQ ID NO: 27.

35. (New): The method defined in claim 28 wherein the nucleic acid comprises SEQ ID NO: 23.

36. (New): The method defined in claim 28 wherein the nucleic acid comprises SEQ ID NO: 25.

37. (New): The method defined in claim 28 wherein the nucleic acid comprises SEQ ID NO: 27.

38. (New): The method defined in claim 31 wherein the nucleic acid comprises SEQ ID NO: 23.

39. (New): The method defined in claim 31 wherein the nucleic acid comprises SEQ ID NO: 25.

40. (New): The method defined in claim 31 wherein the nucleic acid comprises SEQ ID NO: 27.